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NOVEL PEPTIDES FOR TREATMENT OF CANCERFIELD OF INVENTION

This invention relates to novel antiproliferative and anti secretory peptides that are inhibitory to vasoactive intestinal peptide receptor and are useful in the treatment of cancer. The invention particularly relates to the synthesis of lipid-peptide conjugates containing fatty acids of different sizes, which inhibits the binding of VIP to its receptors. The invention encompasses methods for generation of these peptides, composition containing these peptides and the pharmacological applications of these peptides especially in the treatment and prevention of cancer.

BACKGROUND OF THE INVENTION

Vasoactive intestinal peptide (VIP) is a 28-amino acid neuropeptide, which was first isolated from the porcine intestine (Said, S. I. and Mutt, V. , Science, 169, 1217-1218, 1970.) VIP acts as growth factor and plays dominant autocrine and paracrine role in the sustained proliferation of cancer cells. (Said, S.I., Peptides, 5, 143-150, 1984.) Gozes et al. have shown that VIP can serve as autocrine growth factor in lung tumors. (Gozes et al. Biomed. Res. 13 (suppl.2) 37, 1992).

The peptide sequence Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys (SEQ ID NO: 1) is reported to be receptor binding inhibitor of vasoactive intestinal peptide (Said, & Mutt, Ann. N.Y. Acad. Sci., 1, 527, 1988). The role this octapeptide as VIP receptor binding inhibitor has been described in the U.S Patent 5,217,953. In our U.S. Patent Application 08/727,679 we have described the anti cancer role of this VIP binding receptor inhibitor in combination with other neuropeptide analogs. In another U.S. Patent Application 09/248382 we have described the novel analogs of this VIP receptor binding inhibitor incorporating dialkylated amino acids. Keeping in view that lipophilization of bioactive peptides improves their stability, bioavailability and the ability to permeate biomembranes (Dasgupta, P. et al.; 1999, Pharmaceutical Res. 16, 1047-1053; Gozes, I. et al., 1996, Proc. Natl. Acad. Sci. USA, 93, 427-432.), in the present invention we have synthesized lipid conjugates of the peptide sequence Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys (SEQ ID NO: 1) using fatty acids of different sizes, C2- C16 carbon atoms, at the N-terminal site of the peptide. Throughout the application the following abbreviation are used with

the following meanings:

	BOP:	Benzotriazole-1-yl-oxy-tris-(dimethylamino)- phosphonium hexafluorophosphate
5	PyBOP:	Benzotriazole-1-yl-oxy-tris-pyrrolidino-phosphonium hexafluorophosphate
	HBTU:	O-Benzotriazole-N,N,N',N'-tetramethyl-uronium- hexafluoro-phosphate
	TBTU:	2-(1H-Benzotriazole-yl)-1,1,3,3,-tetramethyluronium tetrafluoroborate
10	HOBt:	1-Hydroxy Benzotriazole
	DCC:	Dicyclohexyl carbodiimide
	DIPCD:	Diisopropyl carbodiimide
	DIEA:	Diisopropyl ethylamine
	DMAP:	4-Dimethylamino pyridine
15	DMF:	Dimethyl formamide
	DCM:	Dichloromethane
	NMP:	N-Methyl-2-pyrrolidinone
	TFA:	Trifluoroacetic acid

Throughout the specification and claims, the amino acids residues are
20 designated by their standard abbreviations. Amino acids denote L-configuration
unless otherwise indicated by D or DL appearing before the symbol and separated
from it by hyphen.

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from it by hyphen.

SUMMARY OF THE INVENTION

The present invention relates to peptides of the following general
formula

30 X-Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-Y

wherein,

X is acetyl or straight, branched, or cyclic alkanoyl group of from 3 -

16 carbon atoms.

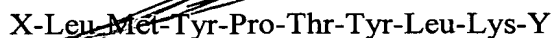
Y is a carboxy terminal residue selected from OH or amino; or a pharmaceutical acceptable salt of the peptides.

BRIEF DESCRIPTION OF THE FIGURE

5 Figure 1 shows the anti-cancer activity of the peptide DT-B1 on PCT xenograft.

DETAILED DESCRIPTION OF THE INVENTION

10 The present invention relates to peptides of the following general formula



wherein,

X is acetyl or straight, branched, or cyclic alkanoyl group of from 3 - 16 carbon atoms.

15 Y is a carboxy terminal residue selected from OH or amino; or a pharmaceutical acceptable salt of the peptides.

The preferred alkanoyl groups are acetyl, n-butanoyl, n-hexanoyl, n-octanoyl, lauroyl, myristoyl, palmitoyl, isohexanoyl, cyclohexanoyl, cyclopentylcarbonyl, n-heptanoyl, n-decanoyl, n-undecanoyl, and 3,7-dimethyloctanoyl.

20 Salts encompassed within the term "pharmaceutically acceptable salts" refer to non-toxic salts of the compounds of this invention. Representative salts and esters include following: acetate, ascorbate, benzenesulfonate, benzoate, bicarbonate, bisulfate, bitartrate, borate, camsylate, carbonate, citrate, dihydrochloride, methanesulfonate, ethanesulfonate, p-toluenesulfonate, 25 cyclohexylsulfamate, quinate, edetate, edisylate, estolate, esylate, fumarate, gluconate, glutamate, glycerophosphates, hydrobromide, hydrochloride, hydroxy-naphthoate, lactate, lactobionate, laurate, malate, maleate, mandelate, mesylate, mucate, napsylate, nitrate, n-methylglucamine, oleate, oxalate, palmoates, pamoate (embonate), palmitate, pantothenate, perchlorates, phosphate/diphosphate, 30 polygalacturonate, salicylates, stearate, succinates, sulfate, sulfamate, subacetate, succinate, tannate, tartrate, tosylate, trifluoroacetate and valerate.

Other salts include Ca, Li, Mg, Na, and K salts; salts of amino acids

such as lysine or arginine; guanidine, diethanolamine or choline; ammonium, substituted ammonium salts or aluminum salts.

The salts are prepared by conventional methods.

The preferred lipo-peptide analogs are:

5 Acetyl-Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-A1) (SEQ ID NO:
2)

n-Butanoyl-Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-B1) (SEQ ID
NO: 3)

n-Octanoyl-Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-01) (SEQ ID
10 NO: 4)

Myristoyl-Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH(DT-M1) (SEQ ID
NO: 5)

Palmitoyl- Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-P1) (SEQ ID
NO: 6)

15 The novel compounds of the present invention have important
pharmacological applications. They are potent anti-neoplastic agents and thereby
possess therapeutic potential in a number of human cancers.

The lipopeptides in the present invention have been generated by
using solid phase techniques or by a combination of solution phase procedures and
20 solid phase techniques or by fragment condensation. The methods for the chemical
synthesis of polypeptides are well known in the art (Stewart and Young, 1969 Solid
Phase Synthesis, W.H. Freeman Co.).

In a preferred embodiment of the present invention the peptides were
synthesized using the Fmoc strategy, on a semi automatic peptide synthesizer (CS
25 Bio, Model 536), using optimum side chain protection. The peptides were
assembled from C-terminus to N-terminus. Peptides amidated at the
carboxy-terminus were synthesized using the Rink Amide resin. The loading of the
first Fmoc protected amino acid was achieved via an amide bond formation with the
solid support, mediated by Diisopropylcarbodiimide (DIPCDI) and HOBt.
30 Substitution levels for automated synthesis were preferably between 0.2 and 0.6
mmole amino acid per gram resin. The steps involved in the synthesis of the
peptide analogs employed the following protocol:

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TABLE I

STEP	REAGENT	MIX TIME (MIN)	NO. OF CYCLES
1.	Methylene chloride	1	2
2.	Dimethyl formamide	1	1
5 3.	20 % Piperidine in Dimethyl formamide	1	1
4.	20 % Piperidine in Dimethyl formamide	29	1
5.	Dimethyl formamide	1	3
6.	Isopropanol	1	2
7.	Methylene chloride	1	2
10 8.	Amino Acid	Variable	1
9.	Dimethyl formamide	1	2
10.	Stop or Return for next cycle		

In a particularly preferred embodiment of the present invention the following chemical moieties were used to protect reactive side chains of the peptides during the synthesis procedure:

The N-terminal amino group was protected by 9-fluorenylmethoxycarbonyl (Fmoc) group. The hydroxyl groups of Threonine and Tyrosine were preferably protected by t-butyl group (tBu). Leu, Met and Pro were used unprotected.

In a preferred embodiment of the invention, 2-8 equivalents of Fmoc protected amino acid per resin nitrogen equivalent were used. The activating reagents used for coupling amino acids to the resin, in solid phase peptide synthesis, are well known in the art. These include DCC, DIPCDI, DIEA, BOP, PyBOP, HBTU, TBTU, and HOBt. Preferably, DCC or DIPCDI / HOBt or HBTU/HOBt and DIEA were used as activating reagents in the coupling reactions. The protected amino acids were either activated in situ or added in the form of preactivated esters known in the art such as N-hydroxy succinamide esters, pentafluorophenyl esters etc. The coupling reaction was carried out in DMF, DCM or NMP or a mixture of these solvents and was monitored by Kaiser test [Kaiser et al., Anal. Biochem., 34,

595-598 (1970)]. In case of a positive Kaiser test, the appropriate amino acid was re-coupled using freshly prepared activated reagents.

After the assembly of the peptide analog was completed, the amino-terminal Fmoc group was removed using steps 1-6 of the above protocol and then the peptide-resin was washed with methanol and dried. The analogs were then deprotected and cleaved from the resin support by treatment with trifluoroacetic acid, crystalline phenol, ethanedithiol, thioanisole and de-ionized water for 1.5 to 5 hours at room temperature. The crude peptide was obtained by precipitation with cold dry ether, filtered, dissolved, and lyophilized.

~~The resulting crude peptide was purified by preparative high performance liquid chromatography (HPLC) using a LiChroCART® C₁₈ (250. Times. 10) reverse phase column (Merck, Darmstadt, Germany) on a Preparative HPLC system (Shimadzu Corporation, Japan) using a gradient of 0.1 % TFA in acetonitrile and water. The eluted fractions were reanalyzed on Analytical HPLC system (Shimadzu Corporation, Japan) using a C₁₈ LiChrospher®, WP-300 (300 X 4) reverse-phase column. Acetonitrile was evaporated and the fractions were lyophilized to obtain the pure peptide. The identity of each peptide was confirmed by electron-spray mass spectroscopy.~~

An analog of the present invention can be made by exclusively solid phase techniques, by partial solid phase/solution phase techniques and/or fragment condensation. Preferred, semi-automated, stepwise solid phase methods for synthesis of peptides of the invention are provided in the examples discussed in the subsequent section of this document.

The present invention will be further described in detail with reference to the following examples, as will be appreciated by a person skilled in the art is merely illustrative and should not be construed as limiting. Various other modifications of the invention will be possible without departing from the spirit and scope of the present invention.

Synthesis of peptides

First loading on Wang Resin

A typical preparation of the Fmoc-Lys-Wang Resin was carried out using 1.0 g of 4-Hydroxymethylphenoxy Resin 1% DVB cross-linked resin (0.7

mM/g) (100-200 mesh), procured from Advanced Chemtech, Louisville, KY, U.S.A. Swelling of the resin was typically carried out in dichloromethane measuring to volumes 10-40ml/g resin. The resin was allowed to swell in methylene chloride (2 X 25 ml, for 10 min.). It was washed once in dimethylformamide (DMF) for 1 min. All solvents in the protocol were added in 20 ml portions per cycle. For loading of the first amino acid on hydroxyl group of the resin, the first amino acid, was weighed in three to six fold excess, along with a similar fold excess of HOBT, in the amino acid vessel of the peptide synthesizer. These were dissolved in dimethylformamide (A.C.S. grade) (J.T.Baker,, New Jersey, U.S.A.) and activated with DIPCDI and 4-dimethyl amino pyridine (DMAP), just prior to the addition to the resin in the reaction vessel of the peptide synthesizer. The coupling reaction was carried out for a period ranging from 6 hours. The loading of the amino acid on the resin was confirmed by the weight gain of the resin. The loading efficiency was ascertained by the increase of weight of the resin after the addition of the amino acid.

EXAMPLE 1

Synthesis of Acetyl-Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-A1)

(SEQ ID NO: 2)

The synthesis of peptide DT-A1 was initiated by using resin loaded with Fmoc-Lys-OH as prepared above on 1g scale. It was subjected to stepwise deprotection and coupling steps as in steps 1-10 of the synthesis cycle. In each coupling reaction, a four-fold excess of appropriate Fmoc amino acid, DIPCDI and HOBT were used. The average coupling time for each amino acids was between 2-5 hrs. On completion of synthesis and removal of the N-terminal Fmoc protecting group (steps 1-6 of the synthesis cycle), the peptideresin was washed twice with methanol. It was further coupled with acetic anhydride in DMF using DIPCDI and HOBT as coupling agents. This was subjected to cleavage in a cleavage mixture consisting of trifluoroacetic acid and scavengers, crystalline phenol, thioanisole, ethanedithol and water for a period of 1-4 hours at room temperature with continuous stirring. The peptide was precipitated using cold dry ether to obtain the crude peptide. The crude peptide was purified on a C₁₈ preparative reverse phase HPLC column (250 X 10) on a gradient system comprising acetonitrile and water in

0.1% TFA as described previously, in the art. The prominent peaks were collected and lyophilized, reanalyzed on analytical HPLC and subjected to mass spectrometry. There was a good agreement between the observed molecular weight and calculated molecular weight (calculated mass = ~ 1070; observed mass = 1071.1). The pure peptide was then used for bioassays.

EXAMPLE 2

Synthesis of n-Butanoyl-Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-B1)

SEQ ID NO: 3

The above peptide sequence was synthesized on resin in a similar way as described in Example 1 except n-butyric anhydride is used in place of acetic anhydride. The final purified peptide was further analyzed by mass spectroscopy. The calculated mass and observed mass was in good agreement (calculated mass = ~ 1098, observed mass = 1099.3).

EXAMPLE 3

Synthesis of n-Octanoyl-Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-O1)

(SEQ ID NO: 4)

The above peptide sequence was synthesized on resin in a similar way as described in Example 1 except octanoic acid is used in place of acetic anhydride. The final purified peptide was further analyzed by mass spectroscopy. The calculated mass is ~ 1154 and observed mass is 1155.2.

EXAMPLE 4

Synthesis of Myristoyl -Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-M1)

(SEQ ID NO: 5)

The above peptide sequence was synthesized on resin in a similar way as described in Example 1 except myristic acid is used in place of acetic anhydride. The final purified peptide was further analyzed by mass spectroscopy (calculated mass = ~ 1238, observed mass = 1239.6).

EXAMPLE 5

Synthesis of Palmitoyl- Leu-Met-Tyr-Pro-Thr-Tyr-Leu-Lys-OH (DT-P1)

(SEQ ID NO: 6)

The above peptide sequence was synthesized on resin in a similar way as described in Example 1 except palmitic acid is used in place of acetic

anhydride. The final purified peptide was further analyzed by mass spectroscopy (calculated mass = ~ 1262, observed mass = 1263.4).

EXAMPLE 6

The cytotoxic effect of Lipo peptide analogs, DT-A1 (SEQ ID NO: 2), DT-B1 (SEQ ID NO: 3), DT-O1 (SEQ ID NO: 4), DT-M1 (SEQ ID NO: 5) and DT-P1 (SEQ ID NO: 6) was studied by MTT assay which is based on the principle of uptake of MTT [3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyl tetrazolium bromide], a tetrazolium salt by the metabolically active cells where it is metabolized by active mitochondria into a blue colored formazan product which can be read spectrophotometrically. Briefly, tumor cells PTC (primary colon) KB (Oral squamous), U87MG (Glioblastoma), HBL100 (Breast), HeP2 (Laryngeal), ECV304 (Endothelial), PA-1 (Ovary) and L132 (Lung) were incubated with the peptide analogs for 48 hours at 37°C in a 96-well culture plate, followed by the addition of 100 µg MTT and further incubation of 1 hour. The formazan crystals formed inside the cells were dissolved with a detergent comprising 10% Sodium dodecyl sulfate and 0.01 N HCl and optical density read on a multiscan ELISA reader. The optical density was directly proportional to the number of proliferating and metabolically active cells. Percent cytotoxicity of peptide analogs is shown in the following Tables.

DT-A1

Cell Line	Percentage cytotoxicity at different concentrations			
	1µM	100n M	10 nM	1nM
KB	21.0 ± 2.3	26.9 ± 2.1	31.0 ± 2.1	25.9 ± 1.6
U87MG	21.9 ± 1.4	26.9 ± 1.5	29.9 ± 2.2	10.4 ± 1.3
HBL100	28.9 ± 1.2	30.5 ± 1.3	31.9 ± 3.5	19.9 ± 3.5
HeP2	19.7 ± 1.1	21.9 ± 2.7	39.9 ± 1.8	14.9 ± 2.2
L132	12.9 ± 2.4	14.6 ± 3.1	26.4 ± 3.2	13.9 ± 2.9
PA-1	6.9 ± 3.2	18.5 ± 2.3	24.9 ± 2.6	21.9 ± 1.5
ECV304	10.8 ± 3.4	22.0 ± 2.3	16.5 ± 3.4	8.0 ± 2.5

DT-B1

Percentage cytotoxicity at different concentrations				
Cell Line	1 μ M	100n M	10 nM	1nM
PTC	31 \pm 1.5	48 \pm 1.7	44 \pm 1.3	36 \pm 1.0
KB	14.8 \pm 2.3	18.9 \pm 3.2	25.9 \pm 4.1	21.9 \pm 0.5
U87MG	20.6 \pm 1.7	30.7 \pm 1.7	39.7 \pm 2.7	33.9 \pm 0.6
HBL100	32.8 \pm 2.2	33.9 \pm 2.8	34.8 \pm 1.8	33.0 \pm 1.5
HeP2	12.9 \pm 4.4	14.9 \pm 5.3	22.9 \pm 1.6	8.7 \pm 1.5
L132	11.8 \pm 2.3	10.7 \pm 2.6	26.8 \pm 1.9	13.9 \pm 1.9
PA-1	33.7 \pm 1.6	38.5 \pm 1.6	45.8 \pm 1.2	43.9 \pm 2.9
ECV304	25.8 \pm 2.3	31.9 \pm 2.8	19.9 \pm 4.2	18.7 \pm 5.3

DT-O1

Percentage cytotoxicity at different concentrations				
Cell Line	1 μ M	100n M	10 nM	1nM
PTC	29 \pm 5.5	33 \pm 3.2	32 \pm 0.3	28 \pm 5.4
KB	23.9 \pm 1.3	26.9 \pm 2.2	34.9 \pm 3.1	21.0 \pm 0.6
U87MG	21.9 \pm 1.5	28.6 \pm 1.2	38.7 \pm 1.7	37.0 \pm 1.6
HBL100	27.4 \pm 2.7	32.8 \pm 2.8	33.7 \pm 2.8	30.0 \pm 1.4
HeP2	18.8 \pm 4.2	17.9 \pm 2.3	22.9 \pm 1.2	8.6 \pm 2.5
L132	7.9 \pm 2.3	14.9 \pm 2.5	25.9 \pm 1.7	19.4 \pm 2.9
PA-1	6.0 \pm 1.4	22.6 \pm 3.6	37.8 \pm 2.2	26.0 \pm 3.9
ECV304	23.9 \pm 2.2	24.9 \pm 2.4	27.9 \pm 3.2	16.9 \pm 1.3

DT-M1

	Percentage cytotoxicity at different concentrations			
Cell Line	1 μ M	100n M	10 nM	1nM
PTC	30 \pm 3.9	31 \pm 4.6	25 \pm 3.6	26 \pm 0.5
KB	12.9 \pm 1.6	19.0 \pm 3.2	23.9 \pm 3.1	21.0 \pm 26
U87MG	5.0 \pm 2.5	32.2 \pm 2.2	44.9 \pm 2.7	29.4 \pm 1.2
HBL100	29.6 \pm 2.3	30.4 \pm 2.4	31.6 \pm 3.8	21.9 \pm 2.4
HeP2	12.9 \pm 1.2	22.9 \pm 1.3	18.7 \pm 2.2	15.8 \pm 2.2
L132	9.6 \pm 2.4	16.8 \pm 2.1	26.7 \pm 1.1	10.6 \pm 1.9
PA-1	15.7 \pm 1.2	25.9 \pm 3.3	42.0 \pm 2.2	27.5 \pm 2.9
ECV304	17.7 \pm 1.2	22.9 \pm 1.0	16.9 \pm 3.1	21.9 \pm 1.6

DT-P1

	Percentage cytotoxicity at different concentrations			
Cell Line	1 μ M	100n M	10 nM	1nM
PTC	31 \pm 3.1	33 \pm 1.1	21 \pm 1.2	27 \pm 3.1
KB	18.0 \pm 2.6	23.0 \pm 2.2	32.0 \pm 2.1	21.9 \pm 2.6
U87MG	18.4 \pm 2.4	32.9 \pm 2.5	34.0 \pm 2.6	9.6 \pm 1.8
HBL100	28.9 \pm 1.3	33.3 \pm 1.4	34.9 \pm 3.6	25.7 \pm 2.5
HeP2	14.9 \pm 1.3	28.9 \pm 1.7	24.9 \pm 2.8	13.9 \pm 1.2
L132	17.8 \pm 2.2	19.6 \pm 1.1	29.0 \pm 1.2	10.9 \pm 1.9
PA-1	21.7 \pm 2.2	25.6 \pm 3.3	21.9 \pm 2.2	20.5 \pm 1.9
ECV304	25.9 \pm 1.4	31.9 \pm 1.3	19.9 \pm 3.4	18.8 \pm 2.6

EXAMPLE 7

In vivo activity of Lipo-peptide analogs

The antitumor activity of DT-B1 (SEQ ID NO: 3) was studied in human colon adenocarcinoma (PTC) xenografts in nude mice. PTC tumor xenografts were grown in Balb/c a thymic mice by subcutaneous inoculation of a single cell suspension of PTC cells (15 X 10⁶ cells/100 μ L). The tumor bearing animals were divided into 2 groups of three animals each including one group

comprising untreated control animals. Treatment with DT-B1 was initiated when the average tumor volumes, as measured using a vernier caliper, were between 1.3 cm³. Solutions of DT-B1 was prepared at a concentration of 126 µg/ml and intravenously administered to the assigned group of tumor bearing animals at a dose of 12.6 µg/100 µL twice a day so that the total dose of 25.2µg/day was administered to each animal. The treatment was continued for a period of 14 days.

The antitumor activity of the compounds was monitored by measuring tumor volumes every fourth day using the formula $W*W*L*0.4$ (W = smaller diameter, L = larger diameter). The percentage inhibition of tumor growth was calculated using the formula $(1 - \text{tumor volume-treated} / \text{tumor volume-control}) * 100$. Figure 1 shows the tumor kinetics till day 20 in the treated and untreated animals. DT-B1 showed a significant antitumor activity on PTC xenografts. The percentage inhibition of tumor growth caused by DT-B1 as compared to controls on day 20 was 95.85%.

All publications referenced are incorporated by reference herein, including the amino acid sequences listed in each publication. All the compounds disclosed and referred to in the publications mentioned above are incorporated by reference herein, including those compounds disclosed and referred to in articles cited by the 10 publications mentioned.

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